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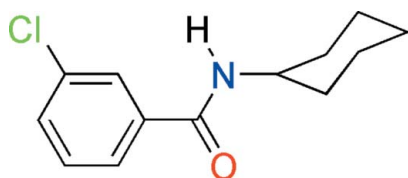
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Key indicators: single-crystal X-ray study; $T = 173$ K; mean $\sigma(\text{C}—\text{C}) = 0.002$ Å; R factor = 0.030; wR factor = 0.069; data-to-parameter ratio = 18.2.

In the title molecule, $\text{C}_{13}\text{H}_{16}\text{ClNO}$, the mean plane of the atoms in the $-\text{CONH}-$ group forms a dihedral angle of $42.0(4)^\circ$ with the benzene ring plane. In the crystal structure, molecules are linked by intermolecular $\text{N}—\text{H} \cdots \text{O}$ hydrogen bonds, generating $C(4)$ chains along $[100]$.

Related literature

For bond-length data, see: Allen (2002). For related structures, see: Garden *et al.* (2005); Wardell *et al.* (2005). For hydrogen-bond motifs, see: Bernstein *et al.* (1995).



Experimental

Crystal data

 $\text{C}_{13}\text{H}_{16}\text{ClNO}$ $M_r = 237.72$ Orthorhombic, $P2_12_12_1$ $a = 8.4963(6)$ Å $b = 11.4891(7)$ Å $c = 12.5318(11)$ Å $V = 1223.29(16)$ Å³ $Z = 4$ Mo $K\alpha$ radiation $\mu = 0.29$ mm⁻¹ $T = 173$ K $0.38 \times 0.22 \times 0.22$ mm

Data collection

Stoe IPDS II two-circle diffractometer

Absorption correction: multi-scan [*MULABS* (Spek, 2003; Blessing, 1995)] $T_{\min} = 0.898$, $T_{\max} = 0.939$

6758 measured reflections

2737 independent reflections

2429 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.035$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.030$ $wR(F^2) = 0.069$ $S = 0.98$

2737 reflections

150 parameters

H atoms treated by a mixture of independent and constrained refinement

 $\Delta\rho_{\max} = 0.18$ e Å⁻³ $\Delta\rho_{\min} = -0.17$ e Å⁻³

Absolute structure: Flack (1983),

1128 Friedel pairs

Flack parameter: 0.03 (5)

Table 1

Hydrogen-bond geometry (Å, °).

$D—H \cdots A$	$D—H$	$H \cdots A$	$D \cdots A$	$D—H \cdots A$
$\text{N1}—\text{H1} \cdots \text{O1}^i$	0.868 (18)	2.052 (18)	2.9161 (15)	173.3 (16)

Symmetry code: (i) $x + \frac{1}{2}, -y + \frac{3}{2}, -z + 1$.

Data collection: *X-Area* (Stoe & Cie, 2001); cell refinement: *X-Area*; data reduction: *X-Area*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL-Plus* (Sheldrick, 2008); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH2817).

References

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supplementary materials

Acta Cryst. (2009). E65, o1265 [doi:10.1107/S1600536809017012]

3-Chloro-*N*-cyclohexylbenzamide

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Comment

We report herein the structure of the title compound, (I) (Fig. 1), which was separated from an impure sample of thiourea by column chromatography as a byproduct, as part of our ongoing studies related to *N,N'*-disubstituted thioureas and heterocyclic compounds. In this class of compounds, N—H···O, C—H···O and N—H···N hydrogen bonds, and weak π – π stacking interactions are the only direction-specific intermolecular interactions (Garden *et al.*, 2005; Wardell *et al.*, 2005). In the crystal structure, molecules form intermolecular N—H···O hydrogen bonds to generate *C*(4) chains (Bernstein *et al.*, 1995) along [100] (Fig. 2). The molecules of (I) are not planar, as evidenced by the torsion angles (C21—N1—C1—O1, 2.9 (02) and C21—N1—C1—C11, -174.88 (11)°) associated with —CONH— moiety, and the amide group adopts the usual *trans* conformation; the bond lengths (Allen, 2002) and inter-bond angles present no unusual values.

Experimental

Freshly prepared and steam distilled 3-chlorobenzoyl isothiocyanate (1.97 g, 10 mmol) was stirred in acetone (30 ml) for 20 min. Neat cyclohexylamin (1.0 g, 10 mmol) was then added and the resulting mixture was stirred for 1 h. The reaction mixture was then poured into 300 ml (approx.) acidified (pH 4) water and stirred well. The solid product was separated and washed with deionized water. One of the fraction obtained during the column chromatography of the target thiourea was recrystallized from methanol/1,1-dichloromethane (1:10 v/v) to give fine crystals of (I), with an overall fractional yield of 15%.

Refinement

H atoms bonded to C were included in calculated positions and refined as riding on their parent C atom with C—H ranging from 0.93 Å to 1.0 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$. The H atom bonded to N was freely refined.

Figures

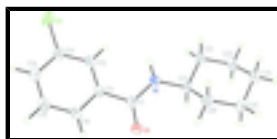


Fig. 1. Molecular structure of (I) showing atom numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.

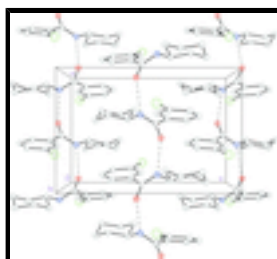


Fig. 2. Part of the crystal structure of (I) viewed onto the *ac* plane. H atoms are omitted for clarity. Dashed lines are drawn between the non-hydrogen donor and acceptor atoms of hydrogen bonds.

3-Chloro-*N*-cyclohexylbenzamide

Crystal data

$C_{13}H_{16}ClNO$	$F_{000} = 504$
$M_r = 237.72$	$D_x = 1.291 \text{ Mg m}^{-3}$
Orthorhombic, $P2_12_12_1$	Mo $K\alpha$ radiation
Hall symbol: P 2ac 2ab	$\lambda = 0.71073 \text{ \AA}$
$a = 8.4963 (6) \text{ \AA}$	Cell parameters from 6652 reflections
$b = 11.4891 (7) \text{ \AA}$	$\theta = 3.4\text{--}27.7^\circ$
$c = 12.5318 (11) \text{ \AA}$	$\mu = 0.29 \text{ mm}^{-1}$
$V = 1223.29 (16) \text{ \AA}^3$	$T = 173 \text{ K}$
$Z = 4$	Block, colourless
	$0.38 \times 0.22 \times 0.22 \text{ mm}$

Data collection

Stoe IPDS II two-circle diffractometer	2737 independent reflections
Radiation source: fine-focus sealed tube	2429 reflections with $I > 2\sigma(I)$
Monochromator: graphite	$R_{\text{int}} = 0.035$
$T = 173 \text{ K}$	$\theta_{\text{max}} = 27.5^\circ$
ω scans	$\theta_{\text{min}} = 3.4^\circ$
Absorption correction: multi-scan [MULABS (Spek, 2003; Blessing, 1995)]	$h = -11 \rightarrow 11$
$T_{\text{min}} = 0.898$, $T_{\text{max}} = 0.939$	$k = -14 \rightarrow 11$
6758 measured reflections	$l = -16 \rightarrow 13$

Refinement

Refinement on F^2	H atoms treated by a mixture of independent and constrained refinement
Least-squares matrix: full	$w = 1/[\sigma^2(F_o^2) + (0.0419P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.030$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.069$	$(\Delta/\sigma)_{\text{max}} = 0.001$
$S = 0.98$	$\Delta\rho_{\text{max}} = 0.18 \text{ e \AA}^{-3}$
2737 reflections	$\Delta\rho_{\text{min}} = -0.17 \text{ e \AA}^{-3}$
150 parameters	Extinction correction: SHELXL97 (Sheldrick, 2008), $F_c^* = kF_c[1 + 0.001 \times F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$
Primary atom site location: structure-invariant direct methods	Extinction coefficient: 0.021 (2)
Secondary atom site location: difference Fourier map	Absolute structure: Flack (1983), 1128 Friedel pairs
Hydrogen site location: inferred from neighbouring sites	Flack parameter: 0.03 (5)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Cl1	0.70767 (6)	1.17100 (4)	0.53982 (4)	0.04673 (13)
C1	0.54914 (14)	0.73267 (12)	0.53979 (11)	0.0203 (3)
O1	0.43572 (10)	0.67397 (9)	0.56950 (7)	0.0245 (2)
N1	0.63351 (13)	0.70808 (10)	0.45221 (9)	0.0233 (2)
H1	0.720 (2)	0.7470 (16)	0.4421 (13)	0.028 (4)*
C11	0.59541 (13)	0.84022 (13)	0.60027 (10)	0.0208 (3)
C12	0.63710 (15)	0.94177 (12)	0.54747 (11)	0.0233 (3)
H12	0.6474	0.9425	0.4720	0.028*
C13	0.66355 (15)	1.04213 (13)	0.60606 (11)	0.0260 (3)
C14	0.65300 (16)	1.04247 (14)	0.71660 (12)	0.0289 (3)
H14	0.6727	1.1117	0.7558	0.035*
C15	0.61351 (17)	0.94075 (15)	0.76869 (12)	0.0300 (3)
H15	0.6069	0.9399	0.8444	0.036*
C16	0.58335 (14)	0.83946 (14)	0.71154 (11)	0.0253 (3)
H16	0.5547	0.7701	0.7479	0.030*
C21	0.59346 (14)	0.61160 (13)	0.38088 (11)	0.0218 (3)
H21	0.4778	0.5966	0.3868	0.026*
C22	0.62972 (17)	0.64688 (13)	0.26615 (11)	0.0274 (3)
H22A	0.7431	0.6654	0.2594	0.033*
H22B	0.5690	0.7176	0.2476	0.033*
C23	0.58727 (17)	0.54859 (16)	0.18917 (12)	0.0334 (4)
H23A	0.4722	0.5353	0.1911	0.040*
H23B	0.6161	0.5717	0.1156	0.040*
C24	0.67181 (18)	0.43660 (15)	0.21786 (13)	0.0340 (3)
H24A	0.7865	0.4471	0.2080	0.041*
H24B	0.6367	0.3735	0.1696	0.041*
C25	0.6385 (2)	0.40235 (15)	0.33291 (14)	0.0373 (4)
H25A	0.7005	0.3322	0.3511	0.045*
H25B	0.5255	0.3827	0.3405	0.045*
C26	0.68006 (17)	0.50066 (13)	0.41084 (12)	0.0292 (3)
H26A	0.6508	0.4774	0.4843	0.035*
H26B	0.7950	0.5148	0.4092	0.035*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.0806 (3)	0.02269 (18)	0.0369 (2)	−0.00841 (19)	−0.0039 (2)	0.00076 (18)
C1	0.0225 (5)	0.0208 (7)	0.0176 (5)	0.0040 (5)	−0.0026 (5)	0.0014 (6)
O1	0.0244 (4)	0.0256 (5)	0.0236 (5)	−0.0026 (4)	0.0023 (3)	0.0014 (4)
N1	0.0236 (5)	0.0234 (6)	0.0227 (5)	−0.0043 (4)	0.0028 (4)	−0.0057 (5)
C11	0.0203 (5)	0.0220 (7)	0.0202 (6)	0.0036 (5)	−0.0005 (4)	−0.0031 (6)
C12	0.0267 (5)	0.0245 (7)	0.0188 (6)	0.0028 (5)	−0.0008 (5)	−0.0020 (6)
C13	0.0304 (6)	0.0213 (7)	0.0261 (7)	0.0022 (5)	−0.0023 (5)	−0.0004 (6)
C14	0.0312 (6)	0.0291 (8)	0.0264 (7)	0.0030 (6)	−0.0031 (5)	−0.0095 (6)
C15	0.0318 (7)	0.0391 (9)	0.0190 (6)	0.0012 (6)	−0.0012 (5)	−0.0058 (6)
C16	0.0273 (6)	0.0292 (8)	0.0195 (6)	0.0012 (5)	0.0002 (5)	0.0004 (6)
C21	0.0231 (6)	0.0224 (7)	0.0199 (6)	−0.0027 (5)	−0.0008 (5)	−0.0054 (5)
C22	0.0355 (7)	0.0253 (7)	0.0214 (6)	0.0044 (6)	−0.0006 (5)	0.0001 (6)
C23	0.0360 (7)	0.0433 (10)	0.0208 (7)	0.0010 (7)	−0.0008 (5)	−0.0076 (7)
C24	0.0391 (8)	0.0306 (8)	0.0322 (7)	−0.0025 (6)	0.0058 (6)	−0.0122 (7)
C25	0.0532 (9)	0.0209 (8)	0.0377 (9)	−0.0028 (7)	0.0057 (7)	−0.0048 (7)
C26	0.0393 (7)	0.0229 (8)	0.0254 (7)	−0.0008 (6)	−0.0016 (6)	0.0018 (6)

Geometric parameters (\AA , $^\circ$)

Cl1—C13	1.7383 (16)	C21—C22	1.5253 (19)
C1—O1	1.2337 (16)	C21—H21	1.0000
C1—N1	1.3410 (17)	C22—C23	1.528 (2)
C1—C11	1.5020 (18)	C22—H22A	0.9900
N1—C21	1.4641 (17)	C22—H22B	0.9900
N1—H1	0.868 (18)	C23—C24	1.517 (3)
C11—C12	1.387 (2)	C23—H23A	0.9900
C11—C16	1.3982 (18)	C23—H23B	0.9900
C12—C13	1.385 (2)	C24—C25	1.521 (2)
C12—H12	0.9500	C24—H24A	0.9900
C13—C14	1.388 (2)	C24—H24B	0.9900
C14—C15	1.380 (2)	C25—C26	1.534 (2)
C14—H14	0.9500	C25—H25A	0.9900
C15—C16	1.390 (2)	C25—H25B	0.9900
C15—H15	0.9500	C26—H26A	0.9900
C16—H16	0.9500	C26—H26B	0.9900
C21—C26	1.519 (2)		
O1—C1—N1	123.32 (13)	C21—C22—C23	110.54 (13)
O1—C1—C11	120.12 (11)	C21—C22—H22A	109.5
N1—C1—C11	116.52 (11)	C23—C22—H22A	109.5
C1—N1—C21	122.34 (11)	C21—C22—H22B	109.5
C1—N1—H1	117.6 (11)	C23—C22—H22B	109.5
C21—N1—H1	119.8 (11)	H22A—C22—H22B	108.1
C12—C11—C16	119.98 (13)	C24—C23—C22	111.43 (12)
C12—C11—C1	121.20 (11)	C24—C23—H23A	109.3

C16—C11—C1	118.62 (13)	C22—C23—H23A	109.3
C13—C12—C11	119.25 (12)	C24—C23—H23B	109.3
C13—C12—H12	120.4	C22—C23—H23B	109.3
C11—C12—H12	120.4	H23A—C23—H23B	108.0
C12—C13—C14	121.38 (14)	C23—C24—C25	110.85 (13)
C12—C13—Cl1	119.39 (11)	C23—C24—H24A	109.5
C14—C13—Cl1	119.22 (12)	C25—C24—H24A	109.5
C15—C14—C13	119.04 (14)	C23—C24—H24B	109.5
C15—C14—H14	120.5	C25—C24—H24B	109.5
C13—C14—H14	120.5	H24A—C24—H24B	108.1
C14—C15—C16	120.66 (13)	C24—C25—C26	111.71 (13)
C14—C15—H15	119.7	C24—C25—H25A	109.3
C16—C15—H15	119.7	C26—C25—H25A	109.3
C15—C16—C11	119.67 (15)	C24—C25—H25B	109.3
C15—C16—H16	120.2	C26—C25—H25B	109.3
C11—C16—H16	120.2	H25A—C25—H25B	107.9
N1—C21—C26	111.83 (11)	C21—C26—C25	110.41 (13)
N1—C21—C22	109.11 (11)	C21—C26—H26A	109.6
C26—C21—C22	110.99 (11)	C25—C26—H26A	109.6
N1—C21—H21	108.3	C21—C26—H26B	109.6
C26—C21—H21	108.3	C25—C26—H26B	109.6
C22—C21—H21	108.3	H26A—C26—H26B	108.1
O1—C1—N1—C21	2.9 (2)	C14—C15—C16—C11	0.9 (2)
C11—C1—N1—C21	−174.88 (11)	C12—C11—C16—C15	−0.08 (19)
O1—C1—C11—C12	−137.24 (12)	C1—C11—C16—C15	−174.93 (12)
N1—C1—C11—C12	40.62 (17)	C1—N1—C21—C26	−92.52 (14)
O1—C1—C11—C16	37.55 (17)	C1—N1—C21—C22	144.31 (12)
N1—C1—C11—C16	−144.60 (12)	N1—C21—C22—C23	−179.27 (11)
C16—C11—C12—C13	−1.17 (19)	C26—C21—C22—C23	57.06 (15)
C1—C11—C12—C13	173.54 (11)	C21—C22—C23—C24	−56.47 (16)
C11—C12—C13—C14	1.6 (2)	C22—C23—C24—C25	55.47 (17)
C11—C12—C13—Cl1	−177.30 (9)	C23—C24—C25—C26	−55.16 (18)
C12—C13—C14—C15	−0.8 (2)	N1—C21—C26—C25	−178.62 (12)
Cl1—C13—C14—C15	178.16 (10)	C22—C21—C26—C25	−56.52 (15)
C13—C14—C15—C16	−0.5 (2)	C24—C25—C26—C21	55.74 (17)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N1—H1 \cdots O1 ⁱ	0.868 (18)	2.052 (18)	2.9161 (15)	173.3 (16)

Symmetry codes: (i) $x+1/2, -y+3/2, -z+1$.

Fig. 1

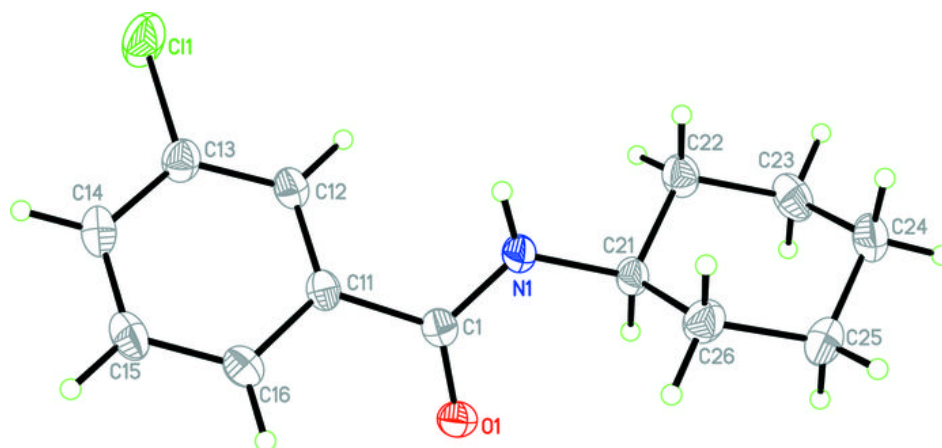


Fig. 2

